

MOLECULAR AND GENOTYPES IDENTIFICATION OF *C. ALBICANS* ISOLATED FROM CHILDREN WITH DIARRHEA IN DIYALA PROVINCE-IRAQ

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ABSTRACT

BACKGROUND

Candida species, one of the most fungi, which isolated from clinical samples that cause many problems to humans, frequently isolated from stools of children with diarrhea.

OBJECTIVE

The aim of the study is, Isolation and identification of *Candida spp.* and genotype of *Candida albicans* infection, among children with diarrhea using conventional methods and PCR, and study distribution of *Candida spp.* and explore the effect of some relevant factors, in Diyala –province.

MATERIALS AND METHODS

A total of 100 children under three years, suffering from diarrhea were included in this study, over a two month period (from 1st October 2016 to 3rd November 2016), in AL-Batool Teaching Hospital. The stool was collected for microscope stool examination, Stool culture on SDA, germ tube test, Chlamydospore formation test, CHROM agar *Candida* (CAC) and PCR techniques.

RESULTS

The results showed that, infection rate by *Candida spp.*, among the patients were (64%), (37.5%) of the total isolates were identified *Candida* as *C. albicans*, (62%) isolates was identified as non-*albicans*., of which (21.9%) isolates were *C. glabrata*, (17.2%) isolate *C. parapsilosis*, (12.5%) isolates was *C. krusei* (10.9%) isolates was *C. tropicalis*. PCR amplification 25S rRNA gene show three genotypes (A, B, C), results showed. (83.3%) for genotype A, (8.3%) for each genotype B and genotype C. the results showed, there are significantly ($P < 0.05$) higher *Candida* infection rate among children with previous antibiotic use and without previous antibiotic, actuality (73.1%), The higher infection rate among children consuming non-sterilization water and children consuming sterilization water, actuality 38(73.1%), Although, insignificant($P > 0.05$), the results showed *Candida* infection rate higher in male patients compared to female patients (66.7% vs. 60.5%) and high *Candida* infection rate in (≥ 2) months patients compared other age groups, actuality 7(77.8%)

CONCLUSIONS

C. albicans is the most common isolated from among the total *Candida* species, and *Candida glabrata* was the most frequent non-*albicans* species. *C. albicans* genotype A, is the most frequent genotype in patients, followed by genotype B and C with the equal rate.

KEYWORDS: *Candida Spp.*, *Candida. Albicans* Genotype & Diarrhea in Children

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INTRODUCTION

Candida species constitute part of the normal microbiota of the human mucosal, oral cavity, vagina, and gastrointestinal tract. Several species, including *Candida albicans*, *C. dublinensis*, *C. glabrata*, *C. guilliermondii*, *C. Lusitaniae*, *C. parapsilosis* and *C. tropicalis*, can be found as part of the normal human commensal flora, especially in all sections of the gastrointestinal tract [1] [2]. In normal, healthy person, there is a balance between *Candida* species, as a normal flora and the normal defense mechanism of the body [3], which will cause opportunistic infection in the presence of any of the predisposing factors like; diabetes mellitus, malnutrition [4], humidity, burn, HIV infection [5], renal failure, endocrine disturbance [6], cancer, indiscriminate usage of antibiotics [7], glucocorticoids and cytotoxic drugs [5]. However, in response to improve or disturbance in the sponsor security systems in the gut, like the intestinal microbiota, gut-associated disease fighting capability, and the mucosal hurdle, *Candida* spp., can convert from safe commercials into pathogens or disturbance in the host defense systems, in the gastrointestinal, including the intestinal microbiota, gut-associated immune system, and the mucosal barrier, *Candida* spp., can convert from harmless commensals into pathogens [8]. Colonization of the gastrointestinal and genitourinary tract, may occur during birth directly from the birth canal [9], at some time during infancy or perhaps later in life, in which, the source may be environmental like polluted fresh and marine water [10], soil, air [11], plant [3], contamination of bedding, hospital wards and wash basins, or could be of human source (mucous membrane or gastrointestinal tract) [12]. *Candida albicans* is the most common *Candida* spp., isolated from human stool. Nevertheless, several reports have suggested that, it may cause diarrhea, as it has also been proposed as a cause of antibiotic-associated diarrhea (AAD), in infants. In recent years, the incidence of *Candida* spp. infections has increased. It has also been shown to be *C. albicans* that also causes diarrhea [13]. Although, not commonly suspected clinically, such pathogenic yeast/yeast-like fungi can increase the severity of diarrhea, causing severe dehydration, malnutrition, and mortality [13].

MATERIAL AND METHODS

SAMPLE COLLECTION

Hundred stool samples were collected from children suffering from diarrhea, during a two month period (from 2nd October 2016 to 3rd December 2016). Less three years attending in AL-Batool Teaching Hospital in Baquba.

CULTURE MEDIA

All the collected samples were inoculated directly, on Sabouraud dextrose agar (SDA) containing Chloramphenicol. The inoculated were kept for incubation, for 48hrs, for Lactophenol cotton blue stain examination of *Candida* spp.

PHENOTYPE IDENTIFICATION OF CANDIDA SPP

Germ tube (GT) test

Rapid diagnostic differentiates *C. albicans* from other species. The inoculums of yeast cells obtained from an isolated colony were suspended in the 0.5 ml of serum, were incubated at 37 °C for 3 hrs. Microscopical examination of the germ tubes, presentation [14].

Drops of CAM were added to the slide and left until becoming dry, a part of four days old colonization has been grown up on SDA; was streaked on the slide, later drops of distilled water were added on the filter paper; to keep it humid and moist. The plate was incubated at 25-35°C for 4-6 days. After an incubation period; a drop of lactophenol blue stain was added to the slide, covered with examined under a microscope (40x) to recognize chlamydospores. [15].

CHROM Agar Candida

Candida was resuscitated by inoculating a loop full of culture from Sabouraud Dextrose Agar into CHROM agar media by streaking a loop full of culture and incubated at 37° for 72hours. After 72-96 hours of incubation, *the Candida colonies* were initially identified by colonial color when compared with standard color photographs supplied by the manufacturer and also presented [16].

IDENTIFICATION GENOTYPE OF *CANDIDA ALBICANS*

DNA Extraction

Isolates of the *Candida albicans* were suspended in 3 ml of (YPD) for 48 hrs., at 37C. Genomic DNA was extracted using Wizard Genomic DNA purification kit (Promega, USA). Extracted DNA was transferred to a sterile 1.5ml Microcentrifuge tube and stored at -20°C.

Primers

The primer pairs used for detection the 25S rRNA were CA-INT-L (5-ATA AGG GAA GTC GGC AAA ATA CCG TAA-3) and CAINT-R (5-CCT TGG CTG TGG TTT CGC TAG ATA GTA GAT-3) as described by McCullough¹³, primers were synthesized by (Bioneer Co., USA). Amplification reactions were performed in the 25µl final volume containing 12.5 master mix (Promega, USA), 1.25µl (25 pmol) each of the primers and 5µl DNA template and complete the volume of PCR grade water. The mixtures reaction was subjected to the following thermal cycling parameters 94oC for 3 min. Followed by 30 cycles of 94oC for 1 min, 55 oC for 1 min, 72oC for 2.5 min and a final extension at 72oC for 10 min following the last cycle. All reaction products by electrophoresis on 1.5% agarose-diamond nucleic acid gel in 1X TBE buffer at 100 V for 1 hrs. And visualized in Gel documentation system. Molecular grade water was included randomly as negative controls and *C. albicans* as a reference in the study.

RESULTS

Culture Media

Morphological culture on Sabouraud dextrose agar SDA medium, the colonies of *C. albicans* colonies on sabouraud dextrose agar were white to creamy, round, soft, and smooth to wrinkled, with a characteristic yeast odor [Figure 1]. Lactophenol cotton blue stain examination of *C. albicans* isolates showed spherical to oval cells, with a presence of budding and was much larger than bacterial cells [Figure 2]. The results of specimen cultures of SDA results showed that, 64 isolates were obtained from 100 samples.



Figure 1: Colonies of *Candida* Spp. Cultured on SDA at 37°C for 48 Hrs (40X)

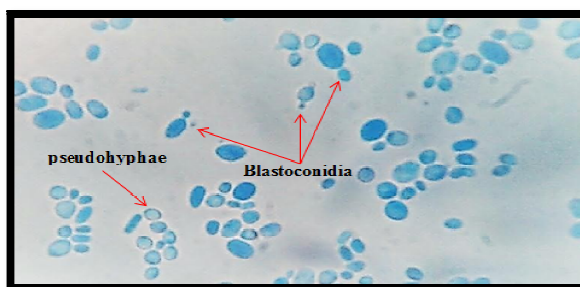


Figure 2: Blastoconidia and Pseudohyphae of the *C. Albicans* Stained With Lactophenol Cotton Blue (40X)

Phenotype Identification of *Candida* Spp

Germ tube (GT) formation was tested, and this test showed that, 24 isolates were positive that the formation of germ tube, which was seen as a long tube-like projections extending from the yeast cells [Figure 3]. Chlamydospore is another test identification character, for identifying *C. albicans* than other species. They were spherical, thick-walled, and usually produced by suppurating cells, that occur along pseudohyphae or at the tip of hyphae. Pseudohyphae and hyphae with clusters of blastospores are also produced, on this agar, [Figure 4]. The colour of colonies on CHROM agar *Candida* was similar, as given by the manufacturer, i. e. green colonies of *C. albicans*. Blue colonies of *C. tropicalis*, Purple- Pink colored colonies of *C. krusei*, *C. glabrata* produced cream to white and *C. parapsilosis* produced pinkish to white, [Figure 5]. The present study shows, difference between *Candida* spp., 24 (37.5%) of the total isolated were identified *Candida*, as *C. albicans*, 40 (62%) isolates were identified as non-*albicans*., of which 14 (21.9%) isolates were *C. glabrata*, 11 (17.2%) isolate *C. parapsilosis* and 8 (12.5%) isolates were *C. krusei* 7 (10.9%) isolates were *C. tropicalis*.



Figure 3: Germ Tube Formation by *C. Albicans* (40X)

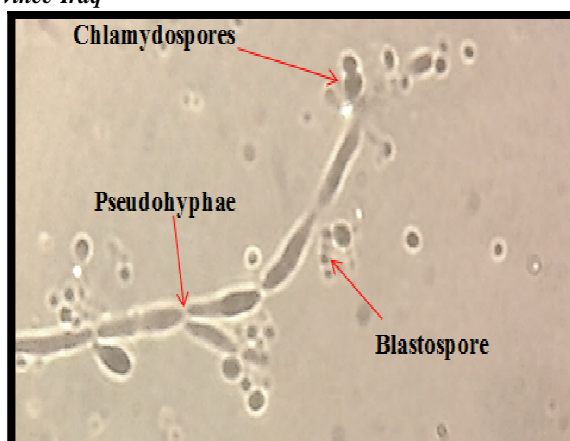


Figure 4: Pseudohyphae, Blastospore and Chlamydospores of *C. Albicans* Cultured on CMA at 30 °C (40X)



Figure 5: colonies of Candida spp A. *C. albicans* B. *c parapsilosis* C, *C.krusei*:D.*C. glabrata* and E.*C. tropicalis* cultured on CHROM agar candida at 37 C for 48 hrs Appeared different colors

Identification Genotype of *Candida Albicans*

Polymerase chain reaction method, using a primer pair designed to span a V3 region in chromosome R, that involve the site of the transposable group I intron of the 25S rRNA gene. Polymerase Chain Reaction amplification, shows three genotypes (A, B, C), results showed single amplification product size (450bp) for 20 (83.3%) isolate, and thus, categorize isolates as genotype A of the *C. albicans*; single amplification product size (840 bp) for 2 (8.3%) isolates and thus, categorize isolates as genotype B *C. albicans* and amplification providing two sizes (450 and 840 bp) for 2 (8.3%) and this categorize isolate as genotype C of the *C. albicans*[Figure 6].

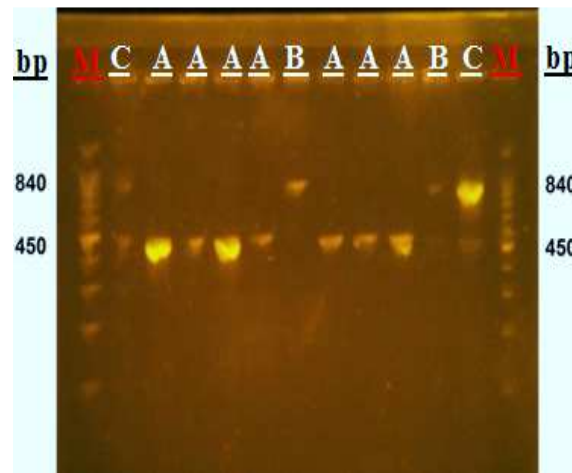


Figure 6: Agarose Gel Electrophoresis of the Candida Albicans Genotype, Stained With Diamond Nucleic Acid Lane M: Molecular Marker (100bp), Lanes A: Genotype A; Lanes B: Genotype B; Lanes C: Genotype C.

In this study, the results showed there are significant ($P < 0.05$) higher *Candida* infection rate, among children with previous antibiotic use and without previous antibiotic, actuality 38 (73.1%) as shown in [Table 1]

Table 1: Candida Infection Rate among Patients According To Previous Antibiotic Use

Previous Antibiotic Use		Total Samples	Candidiasis		Comparison of Significant	
			(+ve)	(-ve)		
Yes	No.	52	38	14	P-value	Sig
	%	100.00%	73.10%	26.90%		
No	No.	48	26	12	0.045	sig.
	%	100.00%	54.20%	45.80%		($P < 0.05$)
Total	No.	100	64	36		
	%	100.00%	64.00%	36.00%		

The highest infection rate among children consuming non-sterilization, water and children consuming sterilized water, actuality 38 (73.1%) as shown in [Table 2]

Table 2: Candida Infection Rate among Patients According to the Water Sterilization

Water Sterilization		Total Samples	Candidiasis		Comparison of Significant	
			(+ve)	(-ve)		
No	No.	52	38	14	P-value	Sig
	%	100.00%	73.1%	26.9%		
Yes	No.	48	26	22	0.045	Sig.
	%	100.00%	54.2	45.8%		($P < 0.05$)
Total	No.	100	64	36		
	%	100.00%	64.00%	36.00%		

Although, insignificant ($P > 0.05$), the results showed a *Candida infection rate* higher in male patients compared to female patients (66.7% vs. 60.5%) as shown in [Table 3], the high *Candida* infection rate in (≥ 2) months patients compared

Table 3: Candida Infection Rate among Patients According to the Gender

Gender		Total Samples	Candidiasis		Comparison of Significant	
			(+ve)	(-ve)		
Male	No.	57	38	19	P-value	Sig
	%	100.00%	66.70%	33.30%		
Female	No.	43	26	17	0.522	Non sig (P>0.05)
	%	100.00%	60.50%	39.50%		
Total	No.	100	64	36		
	%	100.00%	64.00%	36.00%		

Table 4: Candida Infection Rate among Patients According to the Age

Age(Month)		Total Samples	Candidiasis		Comparison of Significant	
			(+Ve)	(-Ve)	P-Value	Sig
≥ 2	No.	9	7	2		
	%	100%	77.80%	22.20%		
3-5	No.	27	14	13	0.585	Non sig. (P>0.05)
	%	100%	51.90%	48.10%		
6-11	No.	40	27	13		
	%	100%	67.50%	32.50%		
12-23	No.	14	9	5		
	%	100%	14.30%	35.70%		
24-36	No.	10	7	3		
	%	100%	70.00%	30.00%		
Total	No.	100	64	36		
	%	100.00%	64.00%	36.00%		

DISCUSSIONS

This study considers as the first report of genotypic analysis of *Candida albicans*, isolated from stools of children with diarrhea in Iraq. The results of this study are agreements with Intesare and Ahmed, that the fungal organism isolated most frequently was *Candida non albican* isolated (62.5%), followed by *Candida albican* (37.5%) Isolated was identified as non-albicans spp., of which (34.4%) isolated was *C. tropicalis* and (9.4%) of remaining isolates were *C. kruzei*, *C. globrata* and *C. parapsilosis*[17]. The result showed that, *C. albicans* the most common isolate from clinical samples and these agree with [18] [19] [20]. In Salah Alden, Ashraf found that, the total rate of *Candida albican* infection was (38%) of 508 examined stool specimens during 2007 at health centers in the Dour town [21]. In Kirkuk province, found that the total rate of *C. albican* infection was (25%) of 200 examined stool specimens, during 2003 at the Pediatric Hospital in Kirkuk [22]. In Najaf governorate, Haydar *et al.* (2013) showed that, 21 (95.45%) isolates belonged to the genotype A and 1 (4.54%) isolates belonged to the genotype B of the *C albicans*. [23]. The results also agree with previous studies showed genotype A of the *C albicans*, predominant in clinical samples, while genotype B, less frequency and this very close to previous studies [24] [25]. In China, it was found that, the rate of genotype A, B, and C of the *C. albicans* from children with early childhood, caries and caries-free children were (61.2%), (15.5%) and (23.3%) [26]. In Northern Ireland, it was found that, rate of genotype A

(71.5%); B (9.5%); C (9.5%) and D (9.5%), were obtained from the blood cultures of patients, with blood borne candidates attending the Belfast City Hospital. In addition, it was found that, the rate of genotype A (66.7%); B (16.7%); C (11.1%) and D (5.1%), were obtained from the following regions (throat, groin, rectum, bronchial washings, sputum, catheter urine and endotracheal tube bio film), attending Belfast City Hospital [27].

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